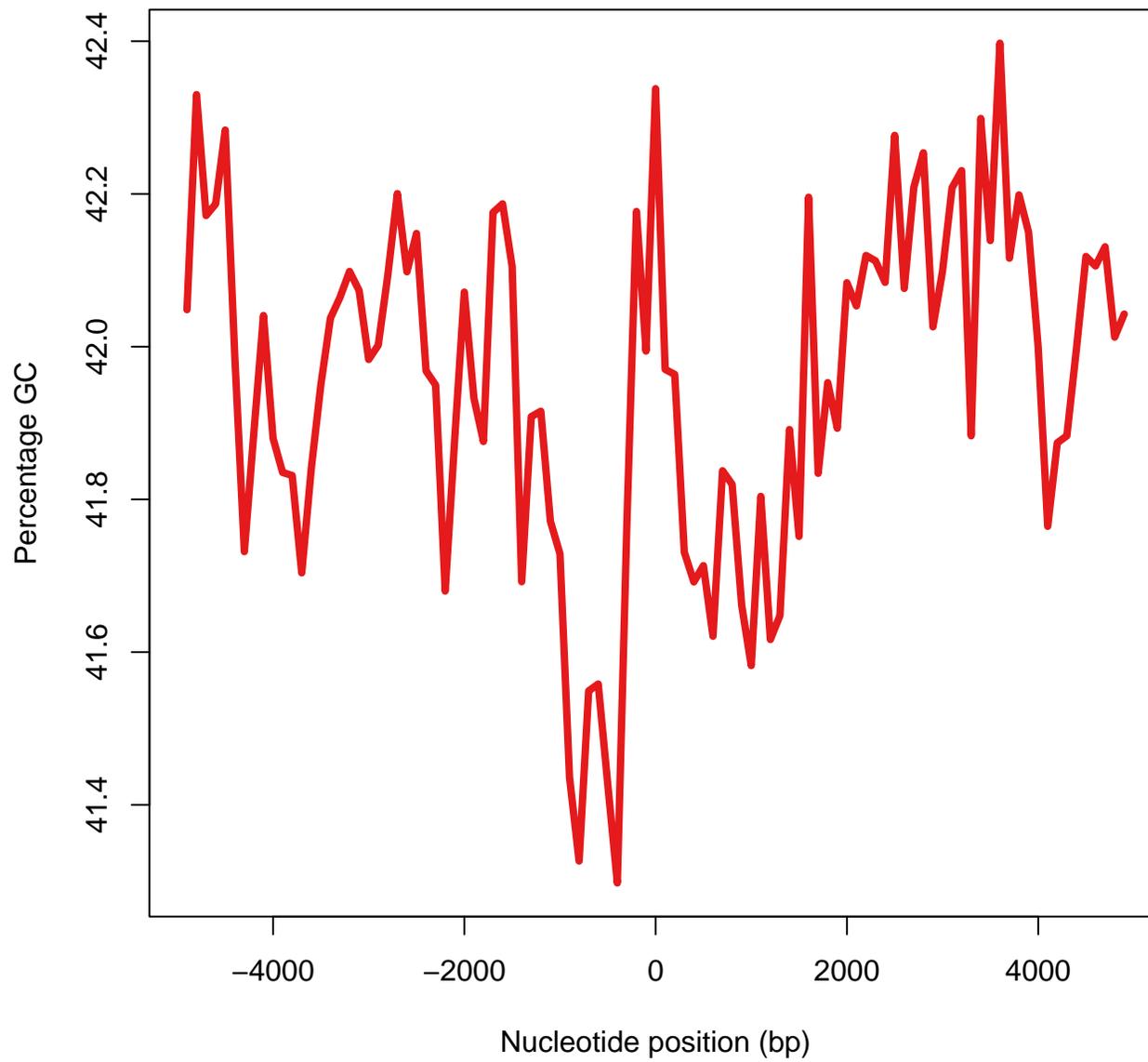
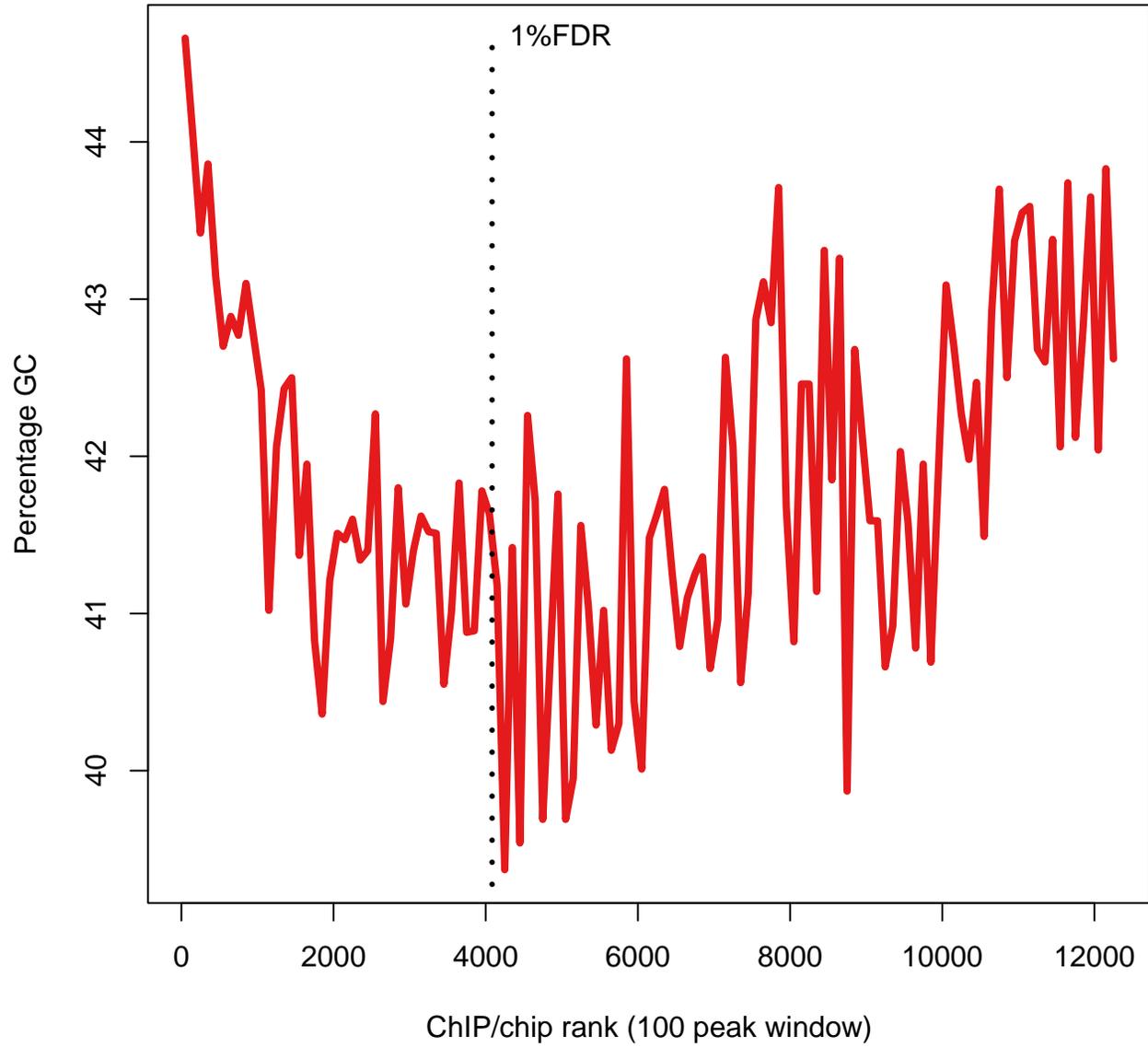


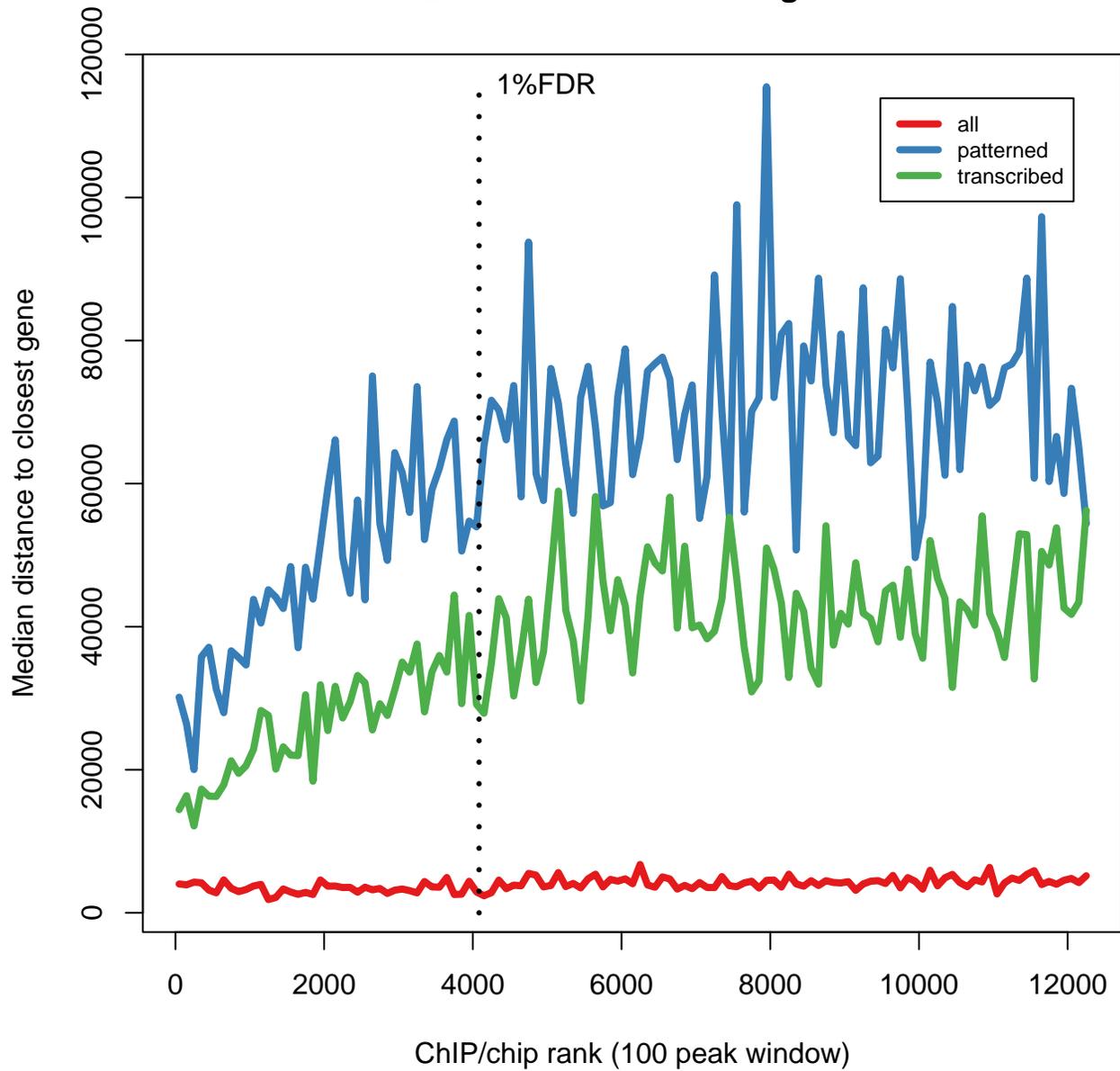
kr_2 base composition across peaks



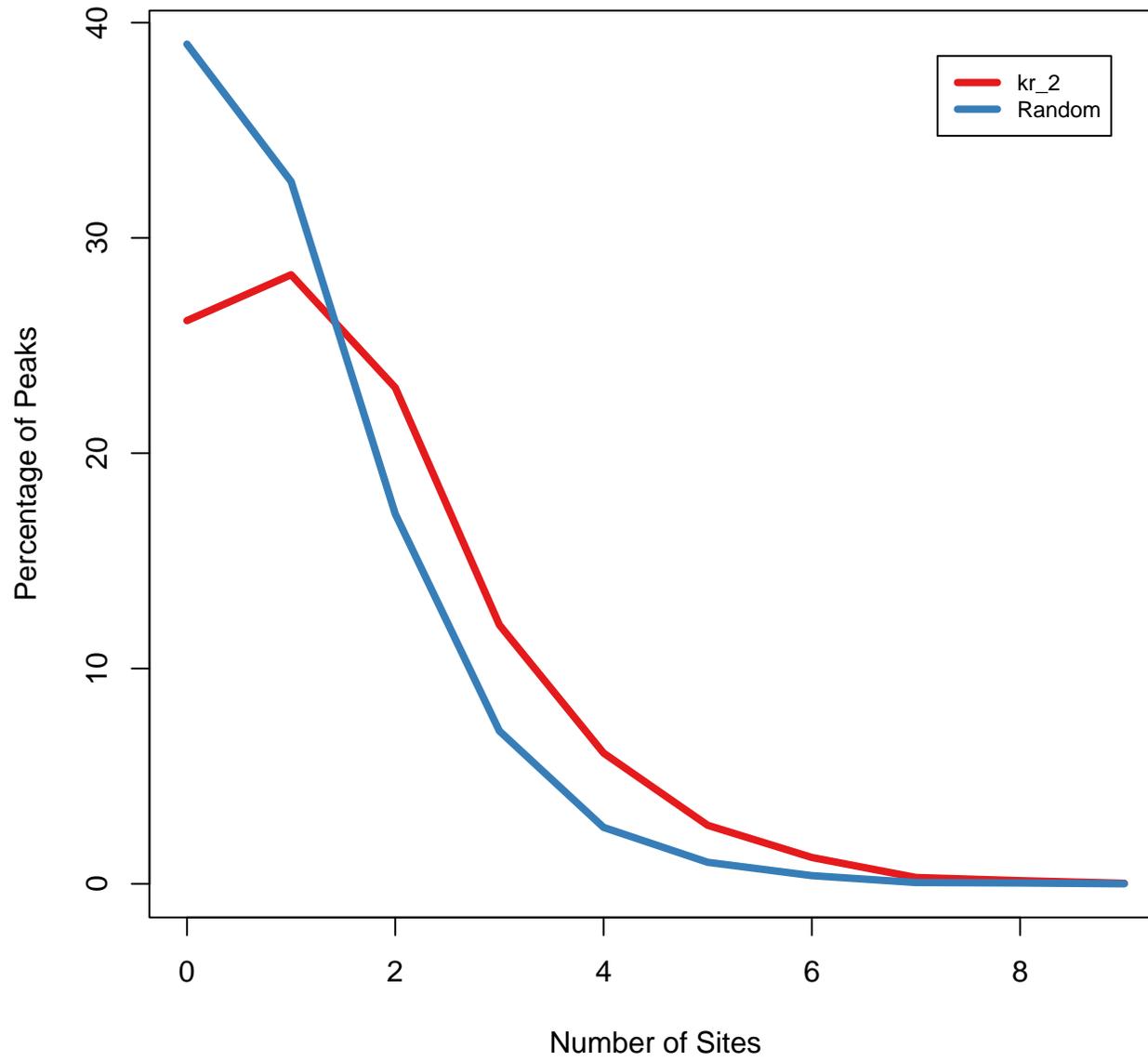
kr_2 base composition down ranks



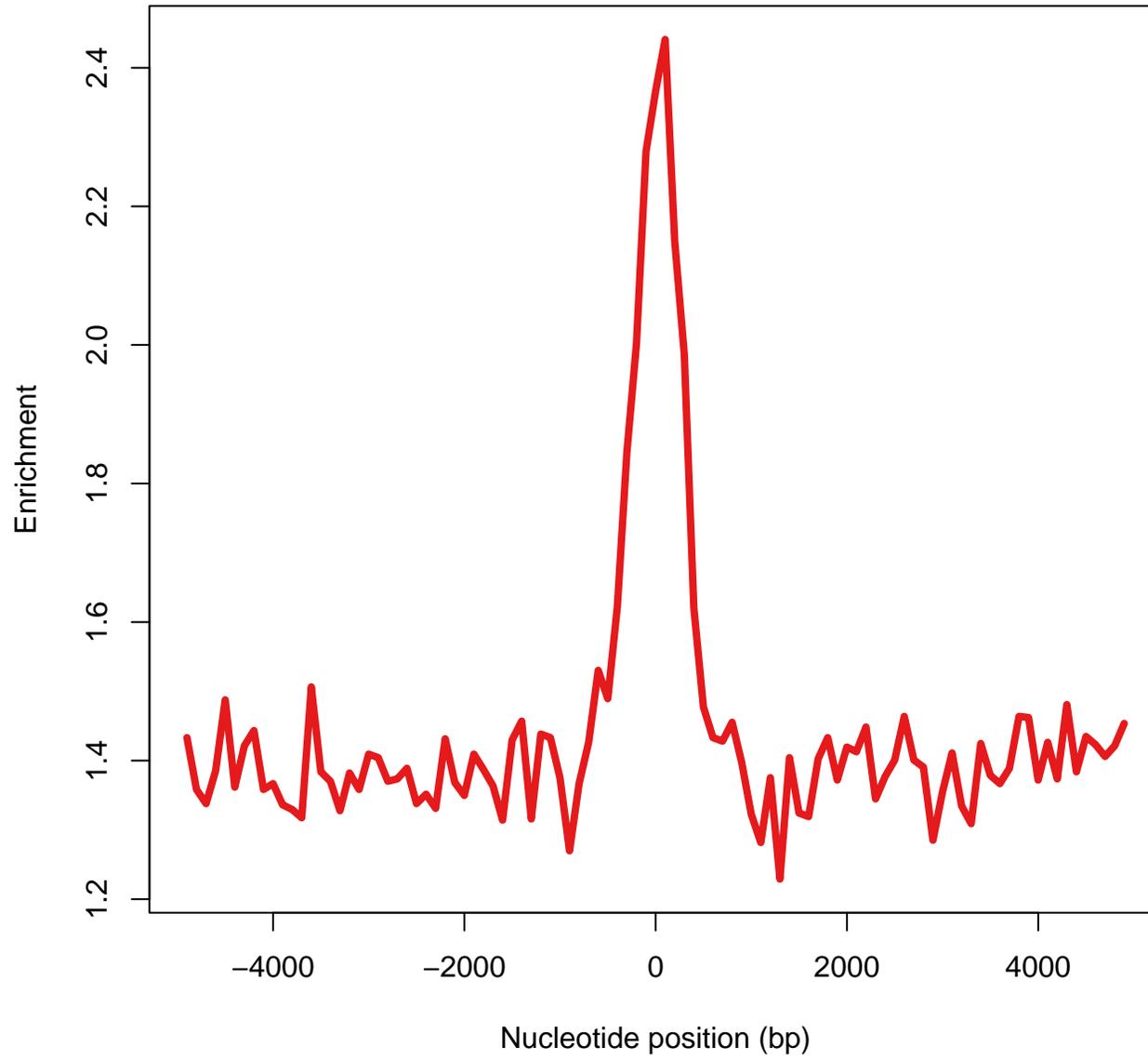
kr_2 median distance to genes



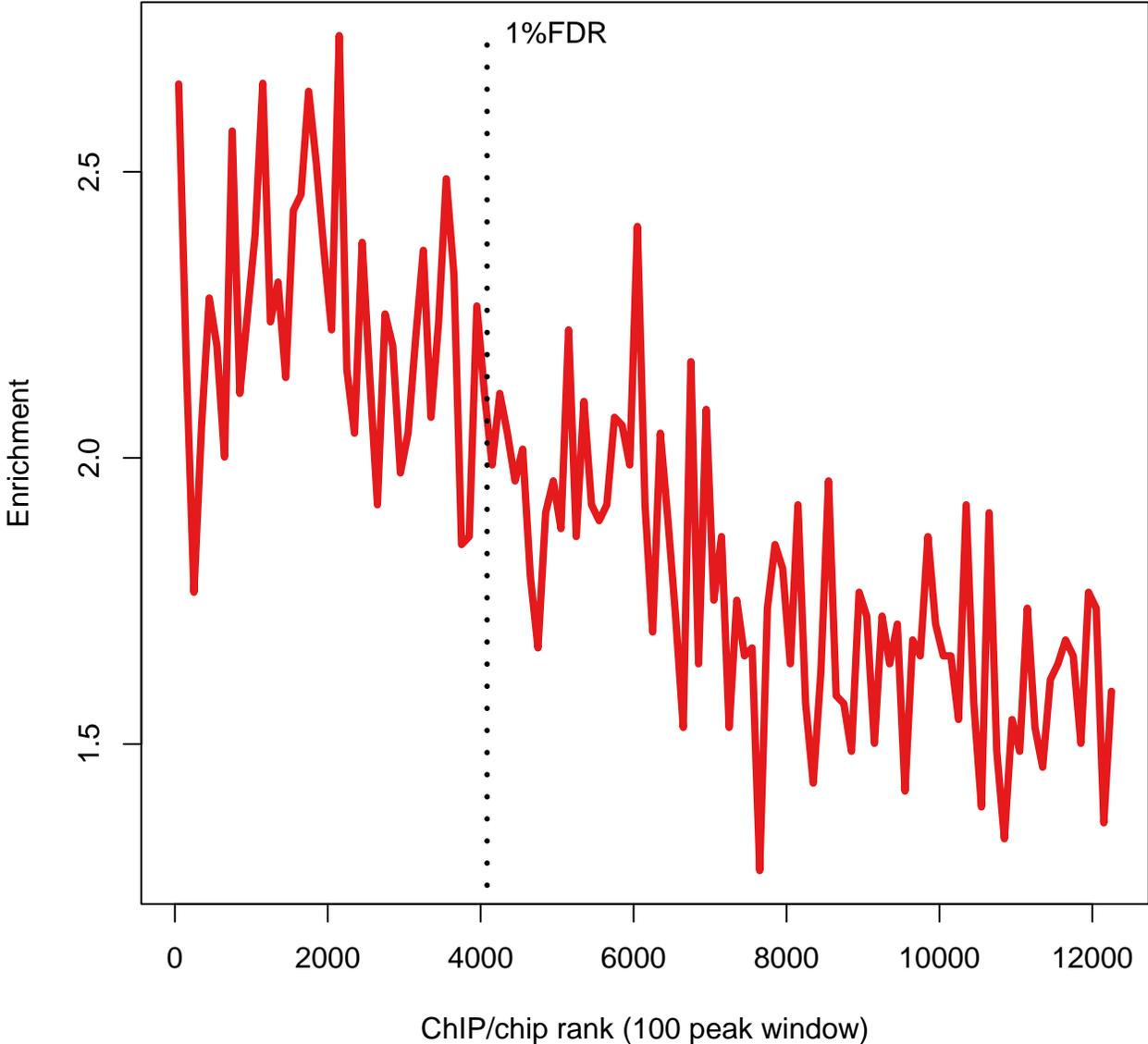
kr_2 distribution of number of sites



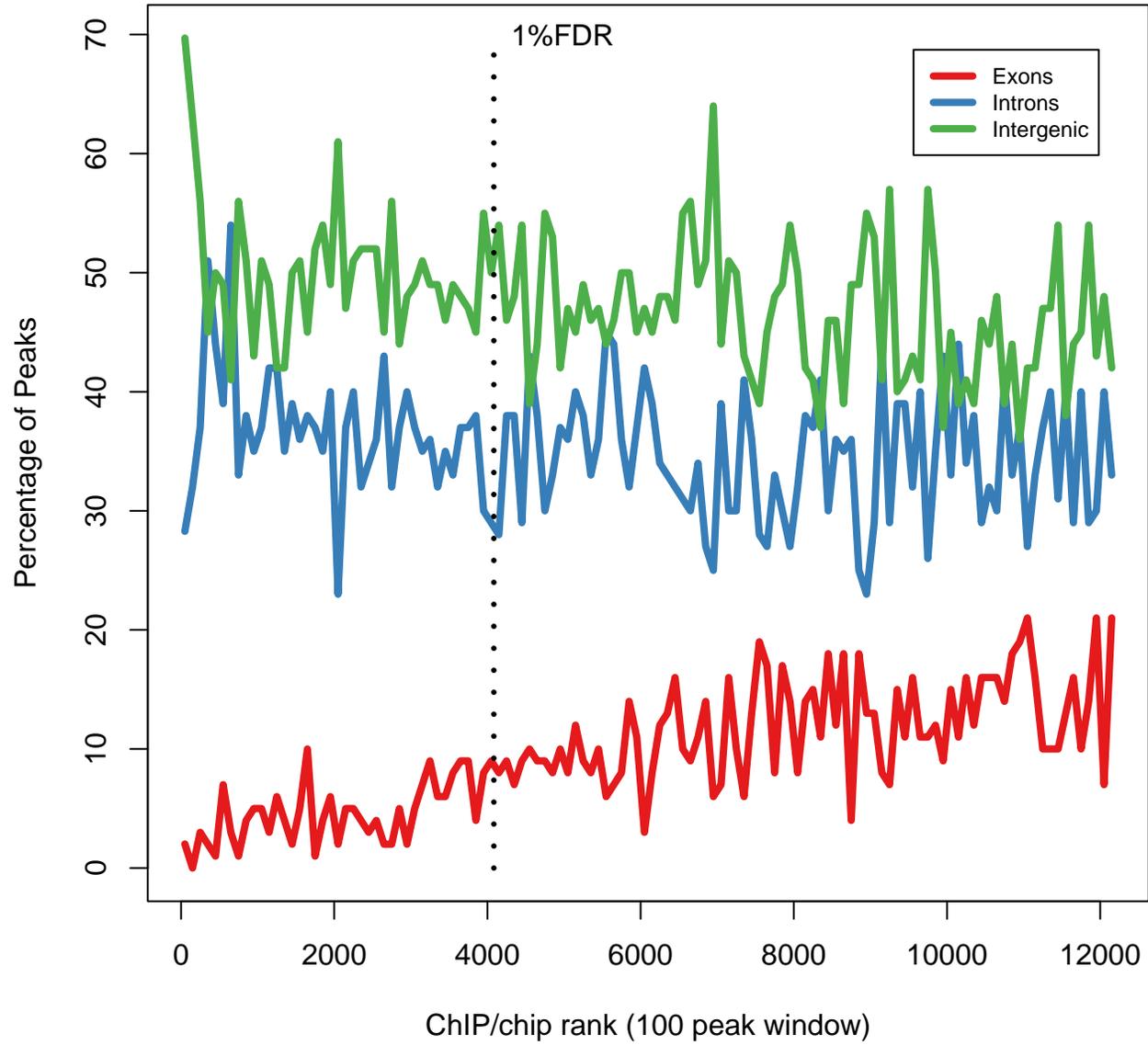
kr_2 PWM enrichment across peaks



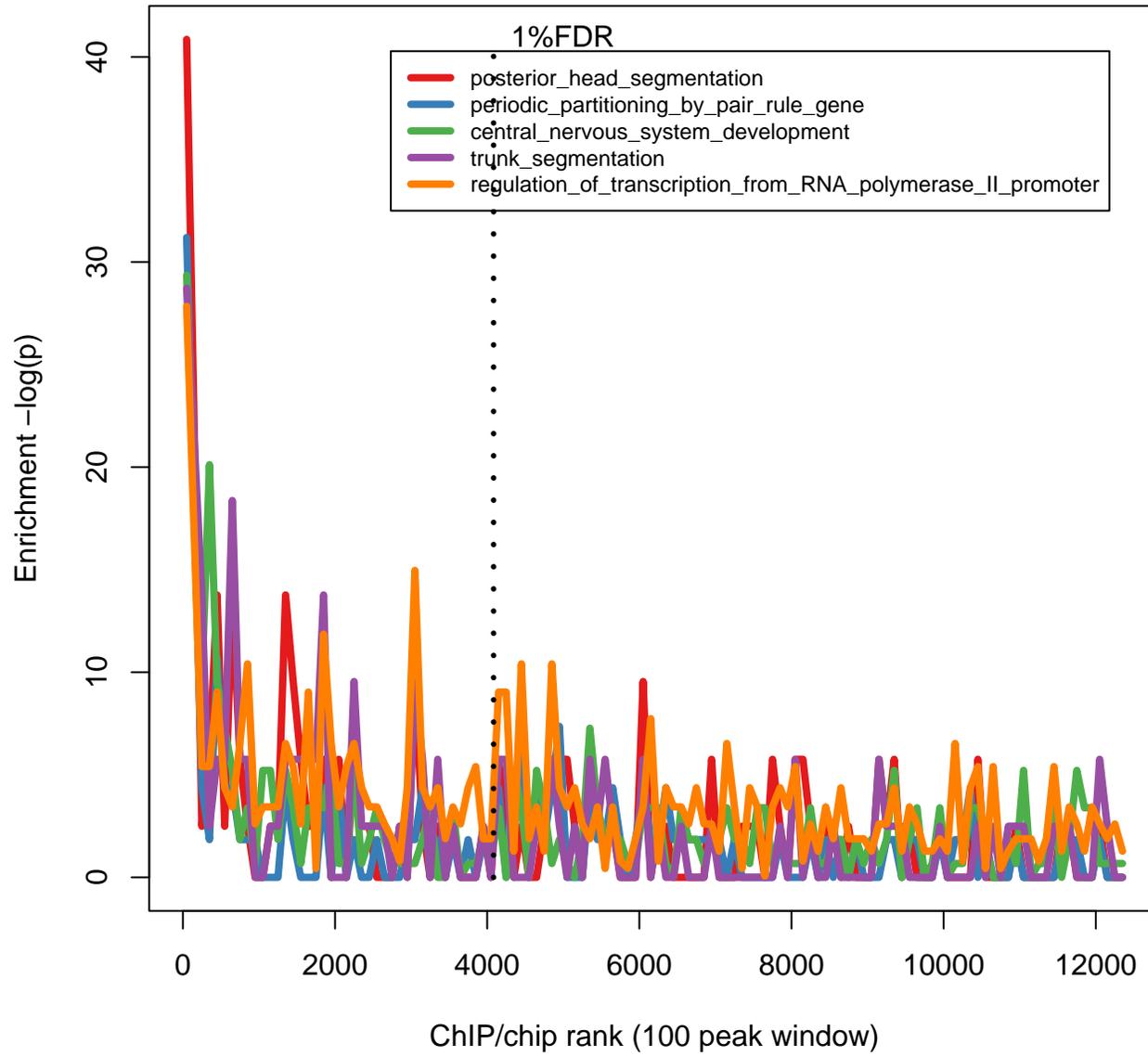
kr_2 PWM enrichment down ranks

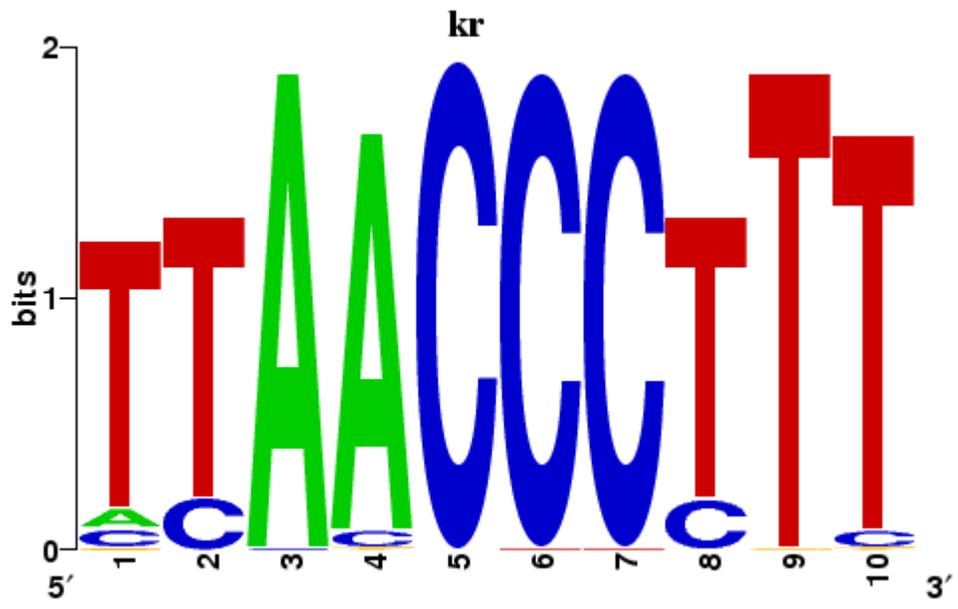


kr_2 genomic location

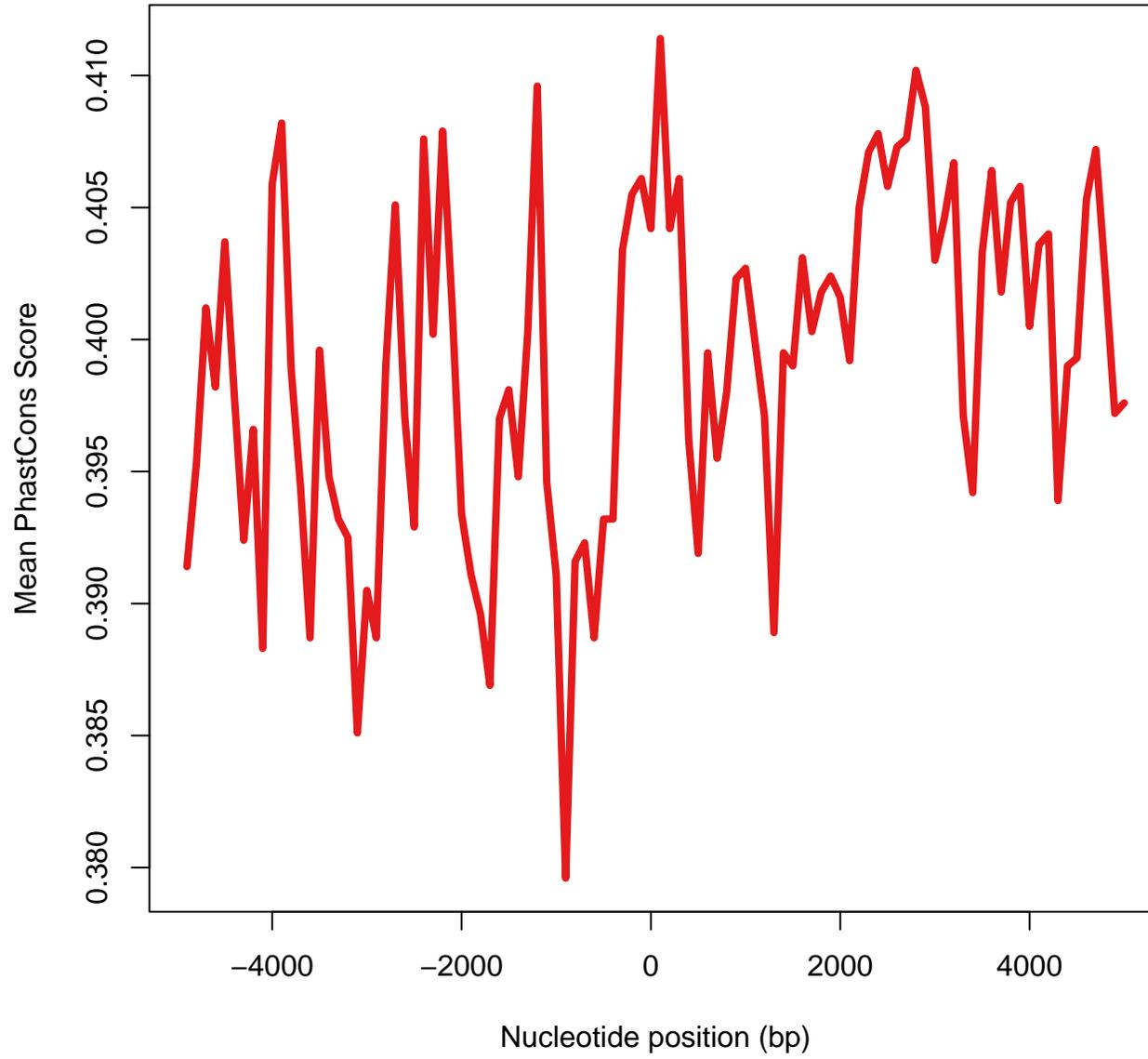


kr_2 GO term enrichment (-logP)





kr_2 mean PhastCons across peaks



kr_2 mean PhastCons down ranks

